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Result
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Maximum Match 10
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Maximum DB seq length: 200000000
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   Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
 protein search, using sw model
                                                                                                                                                                                                                                                                                                                                                                                                                                           100.0
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 PIR_66:*
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258
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Copyright (c) 1993 - 2000 Compugen Ltd
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18.693 Million cell updates/sec
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regulatory protein
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serine/threonine p
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trichohyalin - rab
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homolo
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.,...ernate names: seed storage protein; vicilin precursor C;Species: Gossypium hirsutum (upland cotton) C;Date: 30-Sep 1991 #sequence\_revision 30-Sep-1991 #text\_change 16-Jul-1999 C;Accession: A30838; S06911 R;Chlan, C.A.; Pyle, J.B.; Legocki, A.B.; Dure III, L. Plant Mol. Biol. 7, 475-489, 1986 A;Title: Developmental biochemistry of cottonseed embryogenesis and germinati A;Reference number: A30838 A;Molecule functions.

cottonseed embryogenesis and germination XVIII

RESULT FWCNAB

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YERDPROOYEQCORRCESEATEEREQEQCEORCEREYKEQQROQEEE

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Plant Mol. Biol. 9, 5
A;Title: Developmenta
A;Reference number: S
A;Accession: S06911
                                                                          A; Residues: 1-509 < CHL>C; Superfamily: glycinin
                                                                                                                                                                                                                                                                  alpha-globulin type B precursor (tandem 1) - upland cotton (fragment)
N;Alternate names: seed storage protein
C;Species: Gossypium hirsutum (upland cotton)
C;Date: 31-Mar-1990 #sequence_revision 31-Mar-1990 #text_change 30-Sep-1993
C;Accession: S08059
                                                                                                                                                                                                                                                                                                                                                                           RESULT
S08059
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F;1-24/Domain: signal sequence #status predicted <SIG>
F;25-605/Product: alpha-globulin type A #status predicted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    C:Superfamily: glycinin
C:Keywords: glycoprotein; seed: storage protein
F:1-25/Domain: signal sequence #status predicted <SIG>
F:26-588/Product: alpha-globulin storage proprotein #status predicted
F:417/Binding site: carbohydrate (Asn) (covalent) #status predicted
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A; Residues: 1-81 <CH2>
C:Comment: This is a seed storage protein
                                                                                                    A; Molecule type: DNA
A; Residues: 1-509 < CHL>
                                                                                                                                                                  A; Reference number: S06398
A; Accession: S08059
                                                                                                                                                                                                                              R; Chlan, C.A.; Borroto, K.; Kamalay, J.A.; Plant Mol. Biol. 9, 533-546, 1987
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R;Chlan, C.A.; Borro
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C;Date: 31-Mar-1990 #sequence_revision 31-Mar-1990 #text_change
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A; Accession: S06398
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Query Match
Best Local Similarity
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1. 9, 533-546, 1987
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47.
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Pred. No. 0
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RESULT
S29565
                                                    apolipoprotein A-IV - crab-eating macaque C;Speciles: Macaca fascicularis (crab-eating macaque) C;Date: 06-Jan-1995 #sequence_revision 06-Jan-1995 #C;Accession: S30195; S29565
                    Biochim.
                                   R;Osada,
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                                                                                                                                                                                                                                                                                                                                                                                                                                      J. Biol. Chem. 274, 2563-2570, 1999
A;Title: Multiple functional proteins are produced A;Reference number: Z22767; MUID:99107919
A;Accession: T44430
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              R; Yamada, K.; Shimada, T.; Kondo,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               C; Accession: T44430
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 protein PV100 [imported] - winter squash
C;Species: Cucurbita maxima (winter squash)
C;Date: 21-Jan-2000 #sequence_revision 21-Jan-2000
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Best Local Similarity
Watches 17; Conserv:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  C;Accession: T02634
R;Rieben Jr., W.K.; Gonzales, C.M.; Gonzales, S.T.; Pilkington, K.J.; Kiyosawa, Genetics 148, 1117-1125, 1998
A;Title: Dictyostelium discoldeum nuclear plasmid Ddp5 is a chimera related to the A;Reference number: Z14684; MUID:98198836
A;Accession: T02634
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             rep protein homolog - slime mold (Dictyostelium discoideum) plasmid
C;Species: Dictyostelium discoideum
C;Date: 24-Mar-1999 #sequence_revision 24-Mar-1999 #text_change 29-C
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Best Local Similarity 34.9

Matches 15; Conservative
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                J.; Pocovi, M.; Nicolosi, R.J.; Schaefer, E.J.; Biophys. Acta 1172, 335-339, 1993
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Pred. No.
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apolipoprotein C-III and
                                     Ordovas, J.M
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 A-I
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A;Molecule type: DNA
A;Residues: 1-1898 <LEE>
A;Residues: 1-1898 <LEE>
A;Cross-references: GB:L09190; NID:g292835; PIDN:AAA65582.1; PID:g292836
A;Note: authors translated the codon AGG for residue 1714 as Pro
C;Comment: Trichohyalin is a protein of the medulla of the hair and of the inner root
C;Comment: Trichohyalin is a protein include conversion of arginine to citrulline an
                                                                                                                                                                                                                                                                                                          C:Accession: A45973
R;Lee, S.C.; Kim, I.G.; Marekov, L.N.; O'Reefe, E.J.; Parry, D.A.D.; Steinert, J. Biol. Chem. 268, 12164-12176, 1993
A:Title: The structure of human trichohyalin. Potential multiple roles as a fun ed (cross-linking) protein.
A;Reference number: A45973; MUID:93280194
A;Accession: A45973
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     R;Kimura, M.; Park, S.S.; Sakai, R.; Yamasaki, N.; Funatsu, G. Biosci. Biotechnol. Biochem. 61, 984-988, 1997
A;Title: Primary structure of 6.5k-arginine/glutamate-rich polypeptide from A;Reference number: JC5557; MUID:97357433
A;Accession: JC5557
C;Superfamily: trichohyalin; calmodulin repeat h C;Keywords: calcium binding; citrulline; EF hand F;49-81/Domain: calmodulin repeat homology <EF2>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      δÃ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         C;Comment: This protein is a si F;12-33,16-29/Disulfide bonds:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           C; Species: Luffa cylindrica (smooth roc C; Date: 02-Sep-1997 #sequence_revision
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      arginine/glutamate-rich 6.5K polypeptide - C:Species: Luffa cylindrica (smooth loofah)
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A; Accession: S30195
A; Status: preliminary
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A; Experimental source: seed
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A; Introns: 17/1; 59/2
                                                                         A; Map position:
                                                                                            A; Cross-references: GDB:136223; OMIM:190370
                                                                                                                          A; Gene: GDB: THH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         trichohyalin - human
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A; Residues: 1-429 < OSA>
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                        ;Superfamily: trichohyalin; calmodulin repeat homology;Keywords: calcium binding; citrulline; EF hand; hair;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ;Species: Homo sapiens (man);Date: 03-May_1994 #sequence_revision 01-Mar-1996 #text_change 22-Jun-1999;Date:
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Best Local Similarity 34.8
Matches 16; Conservative
                                                                                                                                             Genetics:
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Best Local
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14; Conservative
                                                                           1921-1921
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s: #status predicted
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Pred. No. 0.21;
6; Mismatches 1
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Pred. No. 0.041;
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A;Reference number: S28589
A;Accession: DNA
A;Residues: 1-1407 <FIE>
A;Cross references: EMBL:Z19092; NID:g1746; PIDN:CAA79519.1; PID:g1747
A;Cross references: EMBL:Z19092; NID:g1746; PIDN:CAA79519.1; PID:g1747
C;Comment: Trichohyalin is a protein of the medulla of the hair and of the inner
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             C:Superfamily: glycinin
():Keywords: seed: storage protein
():Keywords: seed: storage protein
():F:11-20/Domain: signal sequence #status predicted <SIG>
():21-367/Product: prunin 1 alpha chain #status predicted
():368-551/Product: prunin 1 beta chain #status predicted
():368-551/Product: prunin 1 beta chain #status predicted
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                                                                                                                                                                                                                                 submitted to the EMBL Data Library, December 1992 A;Description: Examination of the gene encoding rabbit trichohyalin. A;Reference number: S28589
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A; Residues: 1-60, 'G', 62-551 <GAW>
A; Cross references: EMBL: X78119; I
A; Note: the source is designated
                          A; Introns: 46/3
C; Superfamily:
                                                                                                                                                                                                                                                                                                          C; Accession: S28589
R; Fietz, M.J.; Roge
                                                                                                                                                                                                                                                                                                                                                     C;Species: Oryctolagus cuniculus (domestic rabbit) C;Date: 12-Mar-1993 #sequence_revision 01-Mar-1996
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Plant Mol. Biol. (
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                                                                           C; Genetics:
                                                                                                                                                                                                                                                                                                                                                                                                      trichohyalin -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A; Molecule type: mRNA
A; Residues: 1-551 <GA
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A;Accession: S51941
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Best Local
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                                                  46/3
ly: trichohyalin;
calcium binding;
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18; Conserv
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27, 205-210, 1995
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  calmodulin
citrulline;
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Pred. No. 0.4;
13; Mismatches
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Pred. No. 0.93
9; Mismatches
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  repeat homology
; EF hand; hair;
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  tandem repeat
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RESULT 1.4
T30989
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C;Species: Homo sapiens (man)
C;Jate: 04-Feb-2000 #sequence_revision 04-Feb-2000 #text_change 04-Feb-2000
C;Accession: 746481
R;Duesterhoeft, A; Lauber, J.; Mewes, H.W.; Gassenhuber, J.; Wiemann, S. submitted to the Protein Sequence Database, January 2000
                               serine/threonine protein kinase NIK N;Alternate names: Nck interacting )
                                                                                                                                                                                                                                                                                                                                                                                                                                                          A;Molecule type: mRNA
A;Residues: 1-1027 <AAA>
A;Cross-references: EMBL:AL137755
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       hypothetical protein T20K18.90 - Arabidopsis thaliana
C;Species: Arabidopsis thaliana (mouse-ear cress)
C;Date: 23-Apr-1999 #sequence_revision 23-Apr-1999 #text_change 22-Oct-1999
C;Accession: T06632
R;Bevan, M.; Peters, S.A.; van Staveren, M.; Dirkse, W.; Stiekema, W.; Bancroft, I.; Mew submitted to the Protein Sequence Database, April 1999
A;Reference number: Z15790
A;Accession: T06632
A;Molecule type: DNA
A;Residues: 1-608 <BEV)
A;Residues: 1-608 <BEV)
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A; Accession: T46481
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T46481
                                                                                                                                                                                                                                                                                                                                                                                        A; Note: DKFZp434A025.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A; Status: preliminary
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                В
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A; Introns: 202/3; 216/2; 221/3; 241/3; 292/2; 471/1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A;Gene: ATSP:T20K18.90
A;Map position: 4
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                                                                                                                                                                                                                                                                  Query Match 31.8%; Score 82; DB 2; Best Local Similarity 45.7%; Pred. No. 1.3; Matches 21; Conservative 11; Mismatches 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 31.8%; Score 82; DB 2; Length 608 Best Local Similarity 31.1%; Pred. No. 0.83; Matches 14; Conservative 13; Mismatches 18; Indels
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Best Local
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                                                                                                                                                                             155 EREARROQEREQRRREQE -- EKRRLEELERR -- RKEEEERRRAEEE 196
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                                                                                                                                                                                                                      ERDPROQYEQCORRCESEATEEREQEQCEQRCEREYKEQQRQQEEE 47
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                                    K – mouse
| kinase
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                                                                                                                                                                                                                                                                                                                 Length 1027;
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A;Molecule type: DNA
A;Residues: 1-613 <BES>
A;Cross-references: EMBL:M93690; NID:g159615; PID:g159616
                                                                                                                                                                                                                                                                                                                                                                                                    R;Besansky, N.J.; Paskewitz, S.M.; Mills-Hamm, D.M.; Collins, submitted to the EMBL Data Library, June 1992
A;Description: Distinct families of site-specific retroposons
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         hypothetical protein 1 - African malaria mosquito (fragment) C;Species: Anopheles gambiae (African malaria mosquito) C;Date: 17-Apr-1993 #sequence_revision 17-Apr-1993 #text_cha
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT
S27770
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A:Status: preliminary; translated from GB/EMBL/DDBJ
A:Molecule type: mRNA
A:Residues: 1-1233 <SUY>
A:Cross-references: EMBL:U88984; NID:g1872545; PID:g1872546; PIDN:AAC53165.
C:Keywords: protein kinase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               C:Species: Mus musculus (house mouse)
C:Date: 22-Oct-1999 #sequence_revision 22-Oct-1999 #text_change 22-Oct-1999
C:Accession: T30989
R:Su, Y.C.; Han, J.; Xu, S.; Cobb, M.; Skolnik, E.Y.
EMBO J. 16, 1279-1290, 1997
A:Title: NIK is a new Ste20-related kinase that binds NCK and MEKK1 and actial A:Reference number: Z20954; MUID:97280817
Search co
Job time:
                                                                                                         В
                                                                                                                                                                                                                                                                                                                                                              A; Reference number: S27770
A; Accession: S27770
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R; Besansky, N.J.;
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                                                                                                                                                                                     Query Match 31.4
Best Local Similarity 28.3
Matches 13; Conservative
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Best Local Similarity 45.7
Matches 21; Conservative
                                                                                                  415 EREARROQEREORRREQE--EKRRLEELERR--RKEEEERRRAEEE 456
completed: March
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                                                                                                                                               ERDPRQQYEQCQRRCESEATEEREQEQCEQRCEREYKEQQRQQEEE 47
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                 1,
                                                                                                                                                                                                           31.4%;
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Pred. No.
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                                                                                                                                                                                                                                 2; Length 613;
                                                                                                                                                                                         13;
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